CLAIMS

1. An isolated polynucle otide which encodes a protein comprising the amino acid sequence of SEQ ID NO:2.

2. The isolated polynucleotide of Claim 1, wherein said protein has LysR1 trapscriptional reguatory activity.

3. An isolated polynucleotide, which comprises SEQ ID NO:1.

4. An isolated polynucleotide which is complimentary to the polynucleotide of Claim 3.

5. An isolated polynucleotide which is at least 70% identical to the palynucleotide of Claim 3.

6. An isolated polynuqleotide which is at least 80% identical to the polynucleotide of Claim 3.

7. An isolated polynucled tide which is at least 90% identical to the polynucleotide of Claim 3.

8. An isolated polynucteotide which hybridizes under stringent condition to the polynucleotide of Claim 3; wherein said stringent conditions comprise washing in 5X SSC at a temperature \from 50 to 68°C.

- 9. The isolated polynumleotide of Claim 3, which encodes a protein having LysRi transcriptional regulatory activity.
 - An isolated polynucleotide which comprises at least 15 consecutive nucleotides of the polynucleotide of Claim З.

- The isolated polynucleotide of Claim 10 which comprises SEQ ID NO:3.
- A vector comprising the isolated polynucleotide of Claim 1.

- 13. A vector comprising the isolated polynucleotide of Claim 3.
- 14. A host cell comprising the isolated polynucleotide of Claim 1.
- 5 15. A host cell comprising the isolated polynucleotide of Claim 3.
 - 16. The host cell of Claim 14, which is a Coryneform bacterium.
 - 17. The host cell of Claim 15, which is a Coryneform bacterium.
 - 18. The host cell of Claim 14, wherein said host cell is selected from the group consisting of Coryneform glutamicum, Corynebacterium acetoglutamicum, Corynebacterium acetoacidophilum, Corynebacterium melassecola, Corynebacterium thermoaminogenes, Brevibacterium flavum, Brevibacterium lactofermentum, Brevibacterium divaricatum.
- 19. The host cell of Claim 15, wherein said host cell is selected from the group consisting of Coryneform

 20 glutamicum, Corynebacterium acetoglutamicum,

 Corynebacterium acetoacidophilum, Corynebacterium melassecola, Corynebacterium thermoaminogenes,

 Brevibacterium flavum, Brevibacterium lactofermentum,

 Brevibacterium divaricatum.
- 25 20. A Coryneform bacterium which comprises an attenuated lysR1 gene.
 - 21. The Coryneform bacterium of Claim 21, wherein said lysR1 gene comprises the polynucleotide sequence of SEQ ID NO:1.

. A process for producing L-amino acids comprising culturing a bacterial cell in a medium suitable for producing L-amino acids, wherein said bacterial cell

The process of Claim 23, wherein said bacterial cell

Escherichia Coli DSM 13616.

comprises an attenuated lysR1 gene.

is a Coryneform bacterium or Brevibacterim.

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polynucleotide of Claim 1 to the polynucleotide to be screened; expressing the polynucleotide to produce a protein; and detecting the presence or absence of LysR1 transcriptional regulatory activity in said protein.

- 5 32. A process for screening for polynucleotides which encode a protein having LysR1 transcriptional regulatory activity comprising hybridizing the isolated polynucleotide of Claim 3 to the polynucleotide to be screened; expressing the polynucleotide to produce a protein; and detecting the presence or absence of LysR1 transcriptional regulatory activity in said protein.
 - 33. A method for detecting a nucleic acid with at least 70% homology to nucleotide of Claim 1, comprising contacting a nucleic acid sample with a probe or primer comprising at least 15 consecutive nucleotides of the nucleotide sequence of Claim 1, or at least 15 consecutive nucleotides of the complement thereof.

- 34. A method for producing a nucleic acid with at least 70% homology to nucleotide of Claim 1, comprising contacting a nucleic acid sample with a primer comprising at least 15 consecutive nucleotides of the nucleotide sequence of Claim 1, or at least 15 consecutive nucleotides of the complement thereof.
- 35. A method for detecting a nucleic acid with at least
 70% homology to nucleotide of Claim 3, comprising
 contacting a nucleic acid sample with a probe or primer
 comprising at least 15 consecutive nucleotides of the
 nucleotide sequence of Claim 3, or at least 15
 consecutive nucleotides of the complement thereof.
- 30 36. A method for producing a nucleic acid with at least 70% homology to nucleotide of Claim 3, comprising contacting a nucleic acid sample with a primer comprising at least 15 consecutive nucleotides of the

nucleotide sequence of Claim 3, or at least 15 consecutive nucleotides of the complement thereof.

- 37. A method for making LysR1 protein, comprising: culturing the host cell of Claim 14 for a time and under conditions suitable for expression of LysR1 protein, and collecting the LysR1 protein.
- 38. A method for making LysR1 protein, comprising: culturing the host cell of Claim 15 for a time and under conditions suitable for expression of LysR1 protein, and collecting the LysR1 protein.
- 39. An isolated polypeptide comprising the amino acid sequence of SEQ ID NO:2.

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